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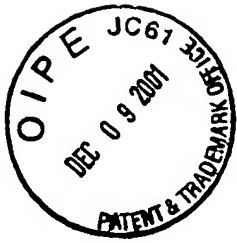


FIGURE 1A

Map of a First IL-17 Receptor Like cDNA (SEQ ID No: 1)  
and Amino Acid (SEQ ID NO: 2)

#5

1 ATAAAAGCGCAGCGTGCGGGTGGCCTGGATCCCGCGCAGTGGCCCGGCGATGTCGCTCGT 60  
M S L V -

61 GCTGCTAAGCCTGGCCGCGCTGTGCAGGAGCGCCGTACCCCGAGAGCCGACCGTTCAATG 120  
L L S L A A L C R S A V P R E P T V Q C -

121 TGGCTCTGAAACTGGGCCATCTCCAGAGTGGATGCTACAACATGATCTAATCCCCGGAGA 180  
G S E T G P S P E W M L Q H D L I P G D -

181 CTTGAGGGACCTCCGAGTAGAACCTGTTACAACACTAGTGTGCAACAGGGGACTATTCAAT 240  
L R D L R V E P V T T S V A T G D Y S I -

241 TTTGATGAATGTAAGCTGGGTACTCCGGGCAGATGCCAGCATCCGCTTGTGTAAGGCCAC 300  
L M N V S W V L R A D A S I R L L K A T -

301 CAAGATTTGTGTGACGGGCAAAAGCAACTTCCAGTCTACAGCTGTGTGAGGTGCAATTA 360  
K I C V T G K S N F Q S Y S C V R C N Y -

361 CACAGAGGCCTTCCAGACTCAGACCAGACCCTCTGGTGGTAAATGGACATTTTCCTACAT 420  
T E A F Q T Q T R P S G G K W T F S Y I -

421 CGGCTTCCCTGTAGAGCTGAACACAGTCTATTTCATTGGGGCCCATAATATTCCTAATGC 480  
G F P V E L N T V Y F I G A H N I P N A -

481 AAATATGAATGAAGATGGCCCTTCCATGTCTGTGAATTTACCTCACCAGGCTGCCTAGA 540  
N M N E D G P S M S V N F T S P G C L D -

541 CCACATAATGAAATATAAAAAAAGTGTGTCAAGGCCGGAAGCCTGTGGGATCCGAACAT 600  
H I M K Y K K K C V K A G S L W D P N I -

601 CACTGCTTGTAAGAAGAATGAGGAGACAGTAGAAGTGAACCTCACAACCACTCCCCTGGG 660  
T A C K K N E E T V E V N F T T T P L G -

661 AAACAGATACATGGCTCTTATCCAACACAGCACTATCATCGGGTTTTCTCAGGTGTTTGA 720  
N R Y M A L I Q H S T I I G F S Q V F E -

721 GCCACACCAGAAGAAACAAACGCGAGCTTCAGTGGTGATTCCAGTGACTGGGGATAGTGA 780  
P H Q K K Q T R A S V V I P V T G D S E -

781 AGGTGCTACGGTGCAGCTGACTCCATATTTTCCTACTTGTGGCAGCGACTGCATCCGACA 840  
G A T V Q L T P Y F P T C G S D C I R H -

841 TAAAGGAACAGTTGTGCTCTGCCCACAAACAGGCGTCCCTTTCCCTCTGGATAACAACAA 900  
K G T V V L C P Q T G V P F P L D N N K -

901 AAGCAAGCCGGGAGGCTGGCTGCCTCTCCTCCTGCTGTCTCTGCTGGTGGCCACATGGGT 960  
S K P G G W L P L L L S L L V A T W V -

Sequence of the first IL-17 Receptor Like cDNA (SEQ ID No: 1) and Amino Acid (SEQ ID NO: 2)

Figure 1B

961	GCTGGTGGCAGGGATCTATCTAATGTGGAGGCACGAAAGGATCAAGAAGACTTCCTTTTC	1020
	L V A G I Y L M W R H E R I K K T S F S -	
1021	TACCACCACACTACTGCCCCCATTAAGGTTCTTGTGGTTTACCCATCTGAAATATGTTT	1080
	T T T L L P P I K V L V V Y P S E I C F -	
1081	CCATCACACAATTTGTTACTTCACTGAATTTCTTCAAACCATTGCAGAAGTGAGGTCAT	1140
	H H T I C Y F T E F L Q N H C R S E V I -	
1141	CCTCGAAAAGTGGCAGAAAAAGAAAATAGCAGAGATGGGTCCAGTGCAGTGGCTTGCCAC	1200
	L E K W Q K K K I A E M G P V Q W L A T -	
1201	TCAAAAAGAGGCAGCAGACAAAGTCGTCTTCCTTCTTTCCAATGACGTCAACAGTGTGTG	1260
	Q K K A A D K V V F L L S N D V N S V C -	
1261	CGATGGTACCTGTGGCAAGAGCGAGGGCAGTCCAGTGAGAACTCTCAAGACCTCTTCCC	1320
	D G T C G K S E G S P S E N S Q D L F P -	
1321	CCTTGCCTTTAACCTTTTCTGCAGTGATCTAAGAAGCCAGATTCATCTGCACAAATACGT	1440
	L A F N L F C S D L R S Q I H L H K Y V -	
1441	GGTGGTCTACTTTAGAGAGATTGATACAAAAGACGATTACAATGCTCTCAGTGTCTGCC	1500
	V V Y F R E I D T K D D Y N A L S V C P -	
1501	CAAGTACCACCTCATGAAGGATGCCACTGCTTTCTGTGCAGAACTTCTCCATGTCAAGCA	1560
	K Y H L M K D A T A F C A E L L H V K Q -	
1561	GCAGGTGTCAGCAGGAAAAAGATCACAAAGCCTGCCACGATGGCTGCTGCTCCTTGTAGCC	1620
	Q V S A G K R S Q A C H D G C C S L *	
1621	CACCCATGAGAAGCAAGAGACCTTAAAGGCTTCCTATCCCACCAATTACAGGGAAAAAC	1680
1681	GTGTGATGATCCTGAAGCTTACTATGCAGCCTACAAACAGCCTTAGTAATTAACATTT	1740
1741	TATACCAATAAAATTTTCAAATATTGCTAACTAATGTAGCATTAACATAACGATTGGAAAC	1800
1801	TACATTTACAACCTCAAAGCTGTTTTATACATAGAAATCAATTACAGCTTTAATTGAAAA	1860
1861	CTGTAACCATTTTGATAATGCAACAATAAAGCATCTTCAGC	1901

**FIGURE 2**  
Homology of a First IL-17 human Receptor Like Polypeptide  
Amino Acid Sequence (SEQ ID NO: 2) and Known Human IL-17  
Receptor Family Member (SEQ ID NO: 3)

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1 .....MSLVLLSLAALCRSAVPREP 20
      || || || |
1 MGAARSPPSAVPGPLLGLLLLLLGV LAPGGASLRLLDHRALVCSQPGLNC 50
21 TVQCGETGPSPPEWMLQHDLPDGLRDLRVEPVTTTSVATGDYSILMNVS 70
      || . | . | . | . | . | . | . | . | . | . | . | . | . | . | . |
51 TVK..NSTCLDDSWIHPRNLTPSSPKDLQIQLFHAHTQQGDLPVVAHIEW 98
71 VLRADASIRLLKATKICVTGKSNFQSYSCVRCNYTEAFQTQTRPSSGGKWT 120
      | . |||| | . . : | . | . ||| . . | . : |
99 TLQTDASILYLEGAELSVLQLNTNERLCVRFE....FLSKLRHHHRRWR 143
121 FSYIGFPVELNTVYFIGAHNIPNANMNEDGPSMSVNFTSPGCLDHIMKYK 170
      | . : | | : . | : | . : | . | . | . | . | . | . |
144 FTFSHFVVDPDQYEYVTVHHLPKPIPDGDPNHQSKNFLVPDCEHARMKVT 193
171 KKC VKAGSLWDPNITACKKNEETVEVNFTTTPLGNRYMALI.....QH 213
      | . . ||||| | . | . | . | . | . | . | . | . | . |
194 TPCMSSGSLWDPNITVETLEAHQLRVSTLWNESTHYQILLTSFPHMENH 243
214 STIIGFSQVFEPHQKQTRASVVIPVTGDSEGA...TVQLTPYFPTCGSD 260
      | . . : | . . . | | . . | . . | . : | . | . | . |
244 SCFEHMHHIPAPRPEEFHQRSNVTLTLRNLKGCCRHQVQIQPFSSCLND 293
261 CIRHKGTVVLCPO.TGVPFPLDNNKSKPGGWLPLLLLSLLVATWVLVAGI 309
      | : || | | | : | : . | . : : | . | . : :
294 CLRHSAT.VSCPEMPDTPPEIPDYMPLWVYWF.ITGISILLVGSVILLIV 341
310 YLMWRHERIKKTSFSTTT.....LLP....PIKVLVVYPSE.ICF 344
      : || | . : | | . | . | . | . | . | . | . : :
342 CMTWRLAGPGSEKYSDDTKYTDGLPAADLIPPPLKPRKVWIIYSADHPLY 391
345 HHTICYFTEFLQNHCRSEVILEKWQKKKIAEMGPVQWLATQK....KAAD 390
      : | : || | . || | : : . | . | . | . | . | .
392 VDVVLKFAQFLLTACGTEVALDLLLEEQAISEAGVMTWVGRQKQEMVESNS 441
391 KVVFLLSNDVNSVCDGTCGKSEGPS.....SENSQDLFPLAFNLCSD 433
      | : : | | . . | . | . | . | . | . | . | . |
442 KIIVLCSRGTRAKWQALLGR..GAPVRLRCDHGKPVGDLFTAAMNMILPD 489
434 LRSQIHLHKYVVVYFREIDTKDDY.NALSVCPKYHLMK..DATAFCAELL 480
      : ||| || | : | . | : | . | : | . | : |
490 FKR PACFGTYVVCYFSEVSCDGDVPDLFGAAPRYPLMDRFEEVYFRIQDL 539
481 HVKQQVSAGKRSQACHDGCCSL*..... 503
      . | . : : |
540 EMFQPGRMHRVGELSGDNYLRSPGGRQLRAALDRFRDWQVRCPDWFECE 589

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FIGURE 3A  
Map of a Second Human IL-17 Receptor Like cDNA (SEQ ID NO: 4)  
And Amino Acid (SEQ ID NO: 5) Sequences

1 ATAAAAGCGC AGCGTGCGGGTG GCCTGGATCCCG CGCAGTGGCCCG GCGATGTCGCTCGT 60  
M S L V -

61 GCTGCTAAG CCTGGCCGCGCT GTGCAGGAGCGC CGTACCCCGAGA GCCGACCGTTCAATG 120  
L L S L A A L C R S A V P R E P T V Q C -

121 TGGCTCTGA AACTGGGCCATC TCCAGAGTGGAT GCTACAACATGA TCTAATCCCCGGA 180  
G S E T G P S P E W M L Q H D L I P G D -

181 CTTGAGGGA CCTCCGAGTAGA ACCTGTTACAAC TAGTGTGCAAC AGGGGACTATTC AAT 240  
L R D L R V E P V T T S V A T G D Y S I -

241 TTTGATGAA TGTAAGCTGGGT ACTCCGGGCAGA TGCCAGCATCCG CTTGTTGAAGGC CAC 300  
L M N V S W V L R A D A S I R L L K A T -

301 CAAGATTTG TGTGACGGGCAA AAGCAACTTCCA GTCCTACAGCTG TGTGAGGCTGGAGTG 360  
K I C V T G K S N F Q S Y S C V R L E C -

361 CAGTGGTGC GATCATGGCTCG CTGCGACCTCAA TCTTCTGGGCTC AAGCGATCGTTC TGC 420  
S G A I M A R C D L N L L G S S D R S A -

421 TTCAGCCTC CCGAGCGGCTGG GACTGCAGGCGT GGGCCACCAGAC CTGGCTAATTTT TGT 480  
S A S R A A G T A G V G H Q T W L I F V -

481 AGTTTTTGT AGAGGGGGGTTT CACCGTGTGCT GGTCTTGAATTC CAGTGCTCAGGC GAT 540  
V F V E G G F T V L L V L N S S A Q A I -

541 CTGCCTGCC TCGGCTTCCCAA AGTGCTGGGATT ACAGTGGACATT TTCCTACATCGG CTT 600  
C L P R L P K V L G L Q W T F S Y I G F -

601 CCCTGTAGA GCTGAACACAGT CTATTTTATTGG GGCCCATATAT TCCTAATGCAAA TAT 660  
P V E L N T V Y F I G A H N I P N A N M -

661 GAATGAAGA TGGCCCTTCCAT GTCTGTGAATTT CACCTCACCAGG CTGCCTAGACCA CAT 720  
N E D G P S M S V N F T S P G C L D H I -

721 AATGAAATA TAAAAAAAAGTG TGTCAAGGCCGG AAGCCTGTGGGA TCCGAACATCAC TGC 780  
M K Y K K K C V K A G S L W D P N I T A -

781 TTGTAAGAA GAATGAGGAGAC AGTAGAAGTGAA CTTACAACCAC TCCCCTGGGAAA CAG 840  
C K K N E E T V E V N F T T T P L G N R -

841 ATACATGGC TCTTATCCAACA CAGCACTATCAT CGGGTTTTCTCA GGTGTTTGAAGCCACA 900  
Y M A L I Q H S T I I G F S Q V F E P H -

901 CCAGAAGAA ACAACGCGAGC TTCAGTGGTGAT TCCAGTGAAGG TGC 960  
Q K K Q T R A S V V I P V T G D S E G A -

961 TACGGTGCA GCTGACTCCATA TTTTCTACTTGG TGGCAGCGACTG CATCCGACATAA AGG 1020  
T V Q L T P Y F P T C G S D C I R H K G -

Figure 3B

1021 AACAGTTGT GCTCTGCCCACA AACAGGCGTCCC TTTCCCTCTGGA TAACAACAAAAG CAA 1080  
T V V L C P Q T G V P F P L D N N K S K -

1081 GCCGGGAGG CTGGCTGCCTCT CCTCCTGCTGTC TCTGCTGGTGGC CACATGGGTGCT GGT 1140  
P G G W L P L L L L S L L V A T W V L V -

1141 GGCAGGGAT CTATCTAATGTG GAGGCACGAAAG GATCAAGAAGAC TTCCTTTTCTAC CAC 1200  
A G I Y L M W R H E R I K K T S F S T T -

1201 CACACTACT GCCCCCATTAA GGTTCCTGTGGT TTACCCATCTGA AATATGTTTCCA TCA 1260  
T L L P P I K V L V V Y P S E I C F H H -

1261 CACAATTTG TTACTTCACTGA ATTTCTTCAAAA CCATTGCAGAAG TGAGGTCATCCT CGA 1320  
T I C Y F T E F L Q N H C R S E V I L E -

1321 AAAGTGGA GAAAAAGAAAAT AGCAGAGATGGG TCCAGTGCAGTG GCTTGCCACTCA AAA 1380  
K W Q K K K I A E M G P V Q W L A T Q K -

1381 GAAGGCAGC AGACAAAGTCGT CTCCTTCTTTC CAATGACGTCAA CAGTGTGTGCGA TGG 1440  
K A A D K V V F L L S N D V N S V C D G -

1441 TACCTGTGG CAAGAGCGAGGG CAGTCCCAGTGA GAACTCTCAAGA CCTCTTCCCCCT TGC 1500  
T C G K S E G S P S E N S Q D L F P L A -

1501 CTTTAACCT TTTCTGCAGTGA TCTAAGAAGCCA GATTCATCTGCA CAAATACGTGGT GGT 1560  
F N L F C S D L R S Q I H L H K Y V V V -

1561 CTACTTTAG AGAGATTGATAC AAAAGACGATTA CAATGCTCTCAG TGTCTGCCCCAA GTA 1620  
Y F R E I D T K D D Y N A L S V C P K Y -

1621 CCACCTCAT GAAGGATGCCAC TGCTTTCTGTGC AGAACTTCTCCA TGTCAAGCAGCA GGT 1680  
H L M K D A T A F C A E L L H V K Q Q V -

1681 GTCAGCAGG AAAAAGATCACA AGCCTGCCACGA TGGCTGTGCTC CTTGTAGCCAC CCA 1740  
S A G K R S Q A C H D G C C S L \*

1741 TGAGAAGCA AGAGACCTTAAA GGCTTCCTATCC CACCAATTACAG GGAAAAAACGTG TGA 1800

1801 TGATCCTGA AGCTTACTATGC AGCCTACAAACA GCCTTAGTAATT AAAACATTTTAT ACC 1860

1861 AATAAAATT TTCAAATATTGC TAACTAATGTAG CATTAACTAACG ATTGGAACTAC ATT 1920

1921 TACAACTTC AAAGCTGTTTTA TACATAGAAATC AATTACAGCTTT AATTGAAACTG TAA 1980

1981 CCATTTTGA TAATGCAACAAT AAAGCATCTTCAGC 2015

GenBank

**FIGURE 4**  
Homology of a Second IL-17 Human Receptor Like Polypeptide  
Amino Acid Sequence (SEQ ID No: 5) and KNOWN Human IL 17  
Receptor Family Mamber (SEQ ID NO: 3)

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1 MSLVLLSLAALCRSAVPREPTVQCGSETGPSPEWMLQHDLPDGLRDLRV 50
1 .....MGAARS 6
51 EPVTTSVATGDYSILMNVSQVLR.ADASIRLL.KATKICVTGKSNFQSYS 98
7 PP..SAVPGPLLGLLLLLLGLVLPAGGASRLRLDHRALVCSQPGLNCTVKN 54
99 CVRLECSGAIMARCDLNLGSSDRSA.....SASRAAGTAGVGHQNWLI 142
55 STCLDDSW.IHPR...NLTPSSPKDLQIQLHFAHTQQGDLPVAHIEWTL 100
143 ....FVVFVEGGFTVLLVLNSSAQAICL..PRLPKVL..GLQWTFYSYIGF 184
101 QTDASILYLEGAELSVLQLNTN.ERLCVRFEFLSKLRHHRRWRFTFSHF 149
185 PVELNTVYFIGAHNIPNANMNEDGPSMSVNFTSPGCLDHIMKYKKKCVKA 234
150 VVDPDQYEYEVTVHHLPKPIPDGDPNHQSKNFLVPDCEHARMKVTTPCMSS 199
235 GSLWDPNITACKKNEETVEVNFTTTPLGNRYMALI.....QHSTIIGF 277
200 GSLWDPNITVETLEAHQLRVSFTLWNESTHYQILLTSFPHMENHSCFEHM 249
278 SQVFEPHQKKQTRASVVIPVTGDSEGA...TVQLTPYFPTCGSDCIRHKG 324
250 HHIPAPRPEEFHQRSNVTLTLRLNLKGCCRQVQIQPFSSCLNDCLRHS 299
325 TVVLC PQ.TGVFPPLDNNKSKPGGWLPLLLLSLLVATWVLVAGIYLMWRH 373
300 T.VSCPEMPDTPEPIPDYMLVWVWF.ITGISILLVGSVILLIVCMTWRL 347
374 ERIKTSFSTTT.....LLP....PIKVLVVYPSE.ICFHHTICY 408
348 AGPGSEKYSDDTKYTDGLPAADLIPPLKPRKVWIIYSADHPLYVDVVLK 397
409 FTEFLQNHCRSEVILEKWQKKKIAEMGPVQWLATQK...KAADKVVFLL 454
398 FAQFLLTACGTEVALDLLEEQAISEAGVMTWVGRQKQEMVESNSKIIVLC 447
455 SNDVNSVCDGTCGKSEGSP.....SENSQDLFPLAFNLFCSDLRSQIH 497
448 SRGTRAKWQALLGR..GAPVRLRCDHGKPVGDLFTAAMNMILPDFKRPAC 495
498 LHKYVVVYFREIDTKDDY.NALSVCPKYHLMK..DATAFCAELLHVKKQV 544
496 FGTYVVCYFSEVSCDGDVPLFGAAPRYPLMDRFEEVYFRIQDLEMFPQG 545
545 SAGKRSQACHDGCCSL*..... 561
546 RMHRVGELSGDNYLRSPGGRQLRAALDRFRDQVRCPDWFECENLYSADD 595

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FIGURE 5A  
Map of a Third IL-17 Receptor Like cDNA (SEQ ID NO: 6)  
and Amino Acid (SEQ ID NO: 7) Sequence

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1 ATAAAGCGCAGCGTGC GGGTGGCCTGGATCCCGCGCAGTGGCCCGGCGATGTCGCTCGT 60
61 GCTGCTAAGCCTGGCCGCGCTGTGCAGGAGCGCCGTACCCCGAGAGCCGACCGTTCAATG 120
121 TGGCTCTGAAACTGGGCCATCTCCAGAGTGGATGCTACAACATGATCTAATCCCGGGAGA 180
181 CTTGAGGGACCTCCGAGTAGAACCTGTTACAACACTAGTGTGCAACAGGGGACTATTCAAT 240
241 TTTGATGAATGTAAGCTGGGTACTCCGGGCAGATGTGGACATTTTCCTACATCGGCTTCC 300
      M W T F S Y I G F P -
301 CTGTAGAGCTGAACACAGTCTATTTTCATTGGGGCCCATAATATTCCTAATGCAAATATGA 360
      V E L N T V Y F I G A H N I P N A N M N -
361 ATGAAGATGGCCCTTCCATGTCTGTGAATTTACCTCACCAGGCTGCCTAGACCACATAA 420
      E D G P S M S V N F T S P G C L D H I M -
421 TGAAATATAAAAAAAAAAGTGTGTCAAGGCCGAAGCCTGTGGGATCCGAACATCACTGCTT 480
      K Y K K K C V K A G S L W D P N I T A C -
481 GTAAGAAGAATGAGGAGACAGTAGAAGTGAAGTTCACAACCACTCCCCTGGGAAACAGAT 540
      K K N E E T V E V N F T T T P L G N R Y -
541 ACATGGCTCTTATCCAACACAGCACTATCATCGGGTTTTCTCAGGTGTTTGAGCCACACC 600
      M A L I Q H S T I I G F S Q V F E P H Q -
601 AGAAGAAACAAACGCGAGCTTCAGTGGTGATTCCAGTGACTGGGGATAGTGAAGGTGCTA 660
      K K Q T R A S V V I P V T G D S E G A T -
661 CGGTGCAGCTGACTCCATATTTTCCTACTTGTGGCAGCGACTGCATCCGACATAAAGGAA 720
      V Q L T P Y F P T C G S D C I R H K G T -
721 CAGTTGTGCTCTGCCCACAAACAGGCGTCCCTTTCCCTCTGGATAACAACAAAAGCAAGC 780
      V V L C P Q T G V P F P L D N N K S K P -
781 CGGGAGGCTGGCTGCTCTCCTCCTGCTGTCTCTGCTGGTGGCCACATGGGTGCTGGTGG 840
      G G W L P L L L L S L L V A T W V L V A -
841 CAGGGATCTATCTAATGTGGAGGCACGAAAGGATCAAGAAGACTTCCTTTTCTACCACCA 900
      G I Y L M W R H E R I K K T S F S T T T -
901 CACTACTGCCCCCATTAAGGTTCTTGTGGTTTACCCATCTGAAATATGTTTCCATCACA 960
      L L P P I K V L V V Y P S E I C F H H T -
961 CAATTTGTTACTTCACTGAATTTCTTCAAAACCATTGCAGAAGTGAGGTCATCCTCGAAA 1020
      I C Y F T E F L Q N H C R S E V I L E K -
1021 AGTGGCAGAAAAAGAAAAATAGCAGAGATGGGTCCAGTGCAGTGGCTTGCCACTCAAAGA 1080
      W Q K K K I A E M G P V Q W L A T Q K K -
1081 AGGCAGCAGACAAAGTCGTCTTCTTCTTTCCAATGACGTCAACAGTGTGTGCGATGGTA 1140
      A A D K V V F L L S N D V N S V C D G T -
1141 CCTGTGGCAAGAGCGAGGGCAGTCCCAGTGAGAACTCTCAAGACCTCTTCCCCCTTGCCT 1200
      C G K S E G S P S E N S Q D L F P L A F -

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Figure 5B

1201 TTAACCTTTTCTGCAGTGATCTAAGAAGCCAGATTTCATCTGCACAAATACGTGGTGGTCT 1260  
N L F C S D L R S Q I H L H K Y V V V Y -

1261 ACTTTAGAGAGATTGATACAAAAGACGATTACAATGCTCTCAGTGTCTGCCCCAAGTACC 1320  
F R E I D T K D D Y N A L S V C P K Y H -

1321 ACCTCATGAAGGATGCCACTGCTTTTCTGTGCAGAACTTCTCCATGTCAAGCAGCAGGTGT 1380  
L M K D A T A F C A E L L H V K Q Q V S -

1381 CAGCAGGAAAAAGATCACAAGCCTGCCACGATGGCTGCTGCTCCTTGTAGCCCACCCATG 1440  
A G K R S Q A C H D G C C S L \*

1441 AGAAGCAAGAGACCTTAAAGGCTTCCTATCCCACCAATTACAGGGAAAAAACGTGTGATG 1500

1501 ATCCTGAAGCTTACTATGCAGCCTACAAACAGCCTTAGTAATTAAAACATTTTATACCAA 1560

1561 TAAAATTTTCAAATATTGCTAACTAATGTAGCATTAAC TAACGATTGGAAACTACATTTA 1620

1621 CAACTTCAAAGCTGTTTTATACATAGAAATCAATTACAGCTTTAATTGAAAAC TGTAAACC 1680

1681 ATTTTGATAATGCAACAATAAAGCATCTTCAGC 1713

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FIGURE 6

Homology of a Third Human IL-17 Receptor Like Polypeptide  
Amino Acid Sequence (SEQ ID NO: 7) and Known Human IL-17  
Receptor Family Member (SEQ ID NO: 3)

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1 .....MWTFSYIGFP 10
101 QTDASILYLEGAELSVLQLNTNERLCVRFEFLSKLRHHHRRWRFTFSHFV 150
11 VELNTVYFIGAHNIPNANMNEDGPSMSVNFTSPGCLDHIMKYKKKCVKAG 60
151 VDPDQYEYEVTVHHLPKPIPDGDPNHQSKNFLVPDCEHARMKVTTPCMSSG 200
61 SLWDPNITACKKNEETVEVNFTTTPLGNRYMALI.....QHSTIIGFS 103
201 SLWDPNITVETLEAHQLRVSFTLWNESTHYQILLTSFPHMENHSCFEHMH 250
104 QVFEPHQKKQTRASVVIPVTGDSEGA...TVQLTPYFPTCGSDCIRHKGT 150
251 HIPAPRPEEFHQRSNVTTLRLNLKGCCRHQVQIQPFSSCLNDCLRHSAT 300
151 VVLCPQ.TGVPPFLDNNKSKPGGWLPLLLLSLLVATWVLVAGIYLMWRHE 199
301 .VSCPEMPDTPEPIPDYMWPLWVYWF.ITGISILLVGSVILLIVCMTWRLA 348
200 RIKKTSFSTTT.....LLP....PIKVLVVYPSE.ICFHHTICYF 234
349 GPGSEKYSDDTKYTDGLPAADLIPPLKPRKVWIIYSADHPLYVDVVLKF 398
235 TEFLQNHRSEVILEKWQKKKIAEMGPVQWLATQK....KAADKVVFLLS 280
399 AQFLLTACGTEVALDLLEEQAISEAGVMTWVGRQKQEMVESNSKIIVLCS 448
281 NDVNSVCDGTCGKSEGSP.....SENSQDLFPLAFNLFCSDLRSQIHL 323
449 RGTRAKWQALLGR..GAPVRLRCDHGKPVGDLFTAAMNMILPDFKRPACF 496
324 HKYVVVYFREIDTKDDY.NALSVCPKYHLMK..DATAFCAELLHVKKQVS 370
497 GTYVVCYFSEVSCDGDVPDLFGAAPRYPLMDRFEEVYFRIQDLEMFQPGR 546
371 AGKRSQACHDGCCSL*..... 386
547 MHRVGELSGDNYLRSPGGRQLRAALDRFRDWQVRCPDWFEENLYSADDQ 596

```

Sequence alignment of IL-17 receptor family members

FIGURE 7  
Overlap of Amino Acid Sequences of the First (SEQ ID NO: 2),  
Second (SEQ ID NO: 5), and Third (SEQ ID NO: 7) Human IL-17  
Receptor Like Polypeptides

1	<b>MSLVLLS LAA</b>	<b>LCRS</b>	<b>AVPREP</b>	<b>TVQCGSETG P</b>	<b>SPEWMLQHDL</b>	<b>IPGDLRDLRV</b>
1	<b>MSLVLLS LAA</b>	<b>LCRS</b>	<b>AVPREP</b>	<b>TVQCGSETG P</b>	<b>SPEWMLQHDL</b>	<b>IPGDLRDLRV</b>
51	<b>EPVTTSVATG</b>	<b>DYSILMNVS</b>	<b>W VLRADASIRL</b>	<b>LKATKICVTG</b>	<b>KSNFQSYSCV</b>	
51	<b>EPVTTSVATG</b>	<b>DYSILMNVS</b>	<b>W VLRADASIRL</b>	<b>LKATKICVTG</b>	<b>KSNFQSYSCV</b>	
101	<b>RCNYTEAFQT</b>	<b>QTRPSGGK</b>	<b>--</b>	<b>-----</b>	<b>-----</b>	<b>-----</b>
101	<b>RLECSGAIMA</b>	<b>RCDLNLGSS</b>	<b>DRSASASRAA</b>	<b>GTAGVGHQNW</b>	<b>LIFVVFVEGG</b>	
119	<b>-----</b>	<b>-----</b>	<b>-----</b>	<b>WTF S</b>	<b>YIGFPVELNT</b>	<b>VYFIGAHNIP</b>
151	<b>FTVLLVLNSS</b>	<b>AQAICLPRLP</b>	<b>KVLGLQWTF S</b>	<b>YIGFPVELNT</b>	<b>VYFIGAHNIP</b>	
1			<b>MWTF S</b>	<b>YIGFPVELNT</b>	<b>VYFIGAHNIP</b>	
143	<b>NANMNEDGPS</b>	<b>MSVNFTSPGC</b>	<b>LDHIMKYKKK</b>	<b>CVKAGSLWDP</b>	<b>NITACKKNEE</b>	
201	<b>NANMNEDGPS</b>	<b>MSVNFTSPGC</b>	<b>LDHIMKYKKK</b>	<b>CVKAGSLWDP</b>	<b>NITACKKNEE</b>	
26	<b>NANMNEDGPS</b>	<b>MSVNFTSPGC</b>	<b>LDHIMKYKKK</b>	<b>CVKAGSLWDP</b>	<b>NITACKKNEE</b>	
193	<b>TVEVNFT TTP</b>	<b>LGNRYMAL IQ</b>	<b>HSTIIGFSQV</b>	<b>FEPHQKKQTR</b>	<b>ASVVIPVTGD</b>	
251	<b>TVEVNFT TTP</b>	<b>LGNRYMAL IQ</b>	<b>HSTIIGFSQV</b>	<b>FEPHQKKQTR</b>	<b>ASVVIPVTGD</b>	
76	<b>TVEVNFT TTP</b>	<b>LGNRYMAL IQ</b>	<b>HSTIIGFSQV</b>	<b>FEPHQKKQTR</b>	<b>ASVVIPVTGD</b>	
243	<b>SEGATVQLTP</b>	<b>YFPTCGSDCI</b>	<b>RHKGTVVLC P</b>	<b>QTGVPPFLDN</b>	<b>NKSKPGGWLP</b>	
301	<b>SEGATVQLTP</b>	<b>YFPTCGSDCI</b>	<b>RHKGTVVLC P</b>	<b>QTGVPPFLDN</b>	<b>NKSKPGGWLP</b>	
126	<b>SEGATVQLTP</b>	<b>YFPTCGSDCI</b>	<b>RHKGTVVLC P</b>	<b>QTGVPPFLDN</b>	<b>NKSKPGGWLP</b>	
293	<u><b>LLLLSLLVAT</b></u>	<u><b>WVLVAGIYLM</b></u>	<u><b>WRHERIKKTS</b></u>	<u><b>FSTTTLLPPI</b></u>	<u><b>KVLVVYPSEI</b></u>	
351	<u><b>LLLLSLLVAT</b></u>	<u><b>WVLVAGIYLM</b></u>	<u><b>WRHERIKKTS</b></u>	<u><b>FSTTTLLPPI</b></u>	<u><b>KVLVVYPSEI</b></u>	
176	<u><b>LLLLSLLVAT</b></u>	<u><b>WVLVAGIYLM</b></u>	<u><b>WRHERIKKTS</b></u>	<u><b>FSTTTLLPPI</b></u>	<u><b>KVLVVYPSEI</b></u>	
343	<b>CFHHTICYFT</b>	<b>EFLQNHCRSE</b>	<b>VILEKWQKKK</b>	<b>IAEMGPVQWL</b>	<b>ATQKKAADKV</b>	
401	<b>CFHHTICYFT</b>	<b>EFLQNHCRSE</b>	<b>VILEKWQKKK</b>	<b>IAEMGPVQWL</b>	<b>ATQKKAADKV</b>	
226	<b>CFHHTICYFT</b>	<b>EFLQNHCRSE</b>	<b>VILEKWQKKK</b>	<b>IAEMGPVQWL</b>	<b>ATQKKAADKV</b>	
393	<b>VFLLSNDVNS</b>	<b>VCDGTCGKSE</b>	<b>GSPSENSQDL</b>	<b>FPLAFNLFCS</b>	<b>DLRSQIHLHK</b>	
451	<b>VFLLSNDVNS</b>	<b>VCDGTCGKSE</b>	<b>GSPSENSQDL</b>	<b>FPLAFNLFCS</b>	<b>DLRSQIHLHK</b>	
276	<b>VFLLSNDVNS</b>	<b>VCDGTCGKSE</b>	<b>GSPSENSQDL</b>	<b>FPLAFNLFCS</b>	<b>DLRSQIHLHK</b>	
443	<b>YVVVYFREID</b>	<b>TKDDYNALSV</b>	<b>CPKYHLMKDA</b>	<b>TAFCAELLHV</b>	<b>KQQVSAGKRS</b>	
501	<b>YVVVYFREID</b>	<b>TKDDYNALSV</b>	<b>CPKYHLMKDA</b>	<b>TAFCAELLHV</b>	<b>KQQVSAGKRS</b>	
326	<b>YVVVYFREID</b>	<b>TKDDYNALSV</b>	<b>CPKYHLMKDA</b>	<b>TAFCAELLHV</b>	<b>KQQVSAGKRS</b>	
493	<b>QACHDGC CSL</b>	<b>*</b>				
551	<b>QACHDGC CSL</b>	<b>*</b>				
376	<b>QACHDGC CSL</b>	<b>*</b>				

Figure 8

# Northern Blot Expression Analysis of TH00-018 Necropsied Transgenic Founders

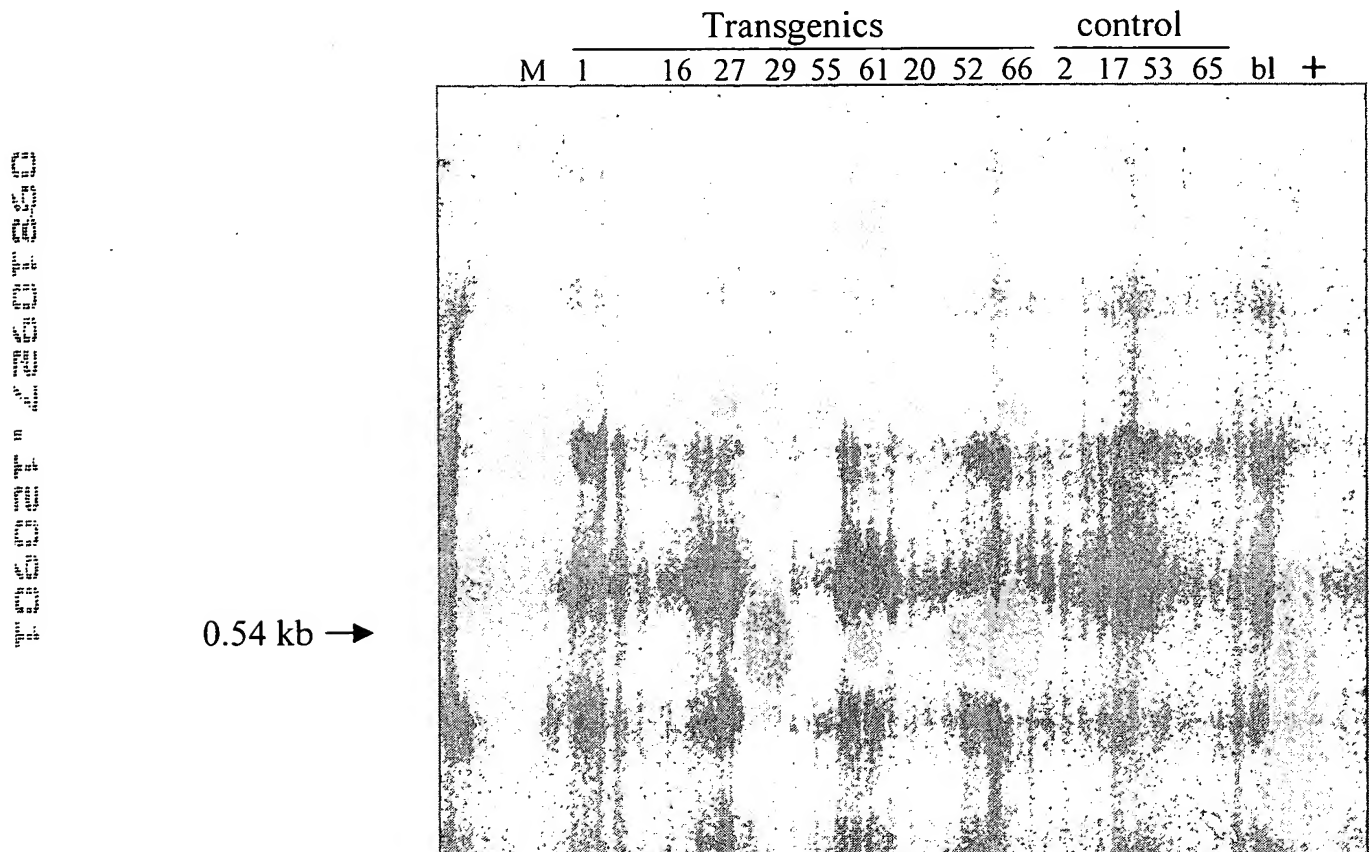


Figure 9

Northern Blot Expression Analysis of TH00-018  
Hepatectomized Transgenic Founders

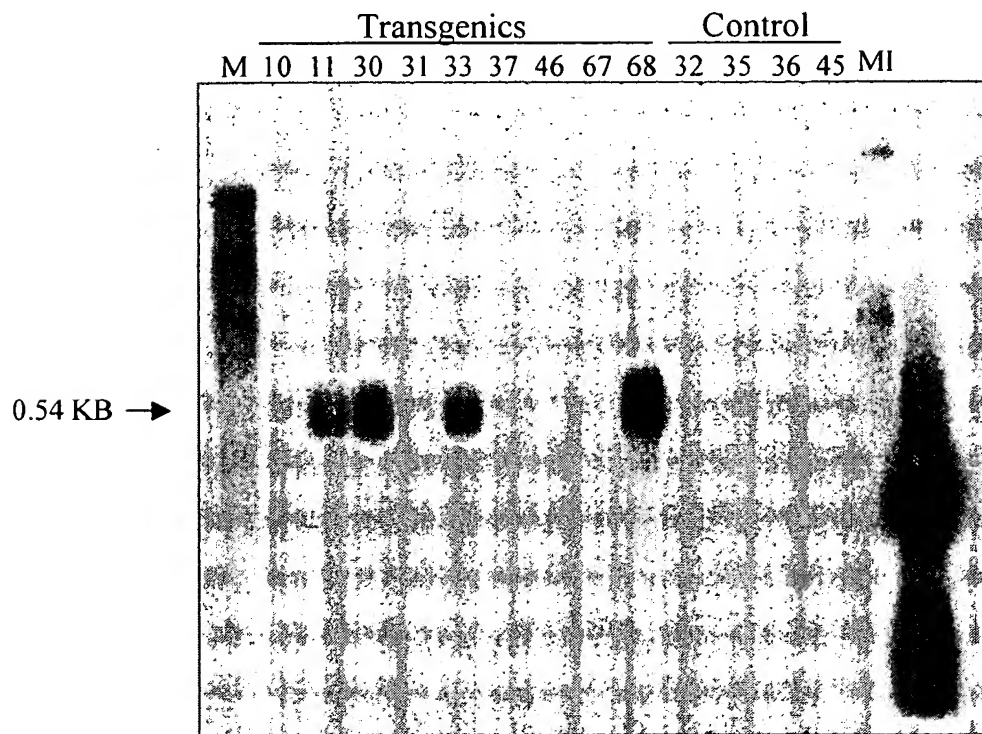
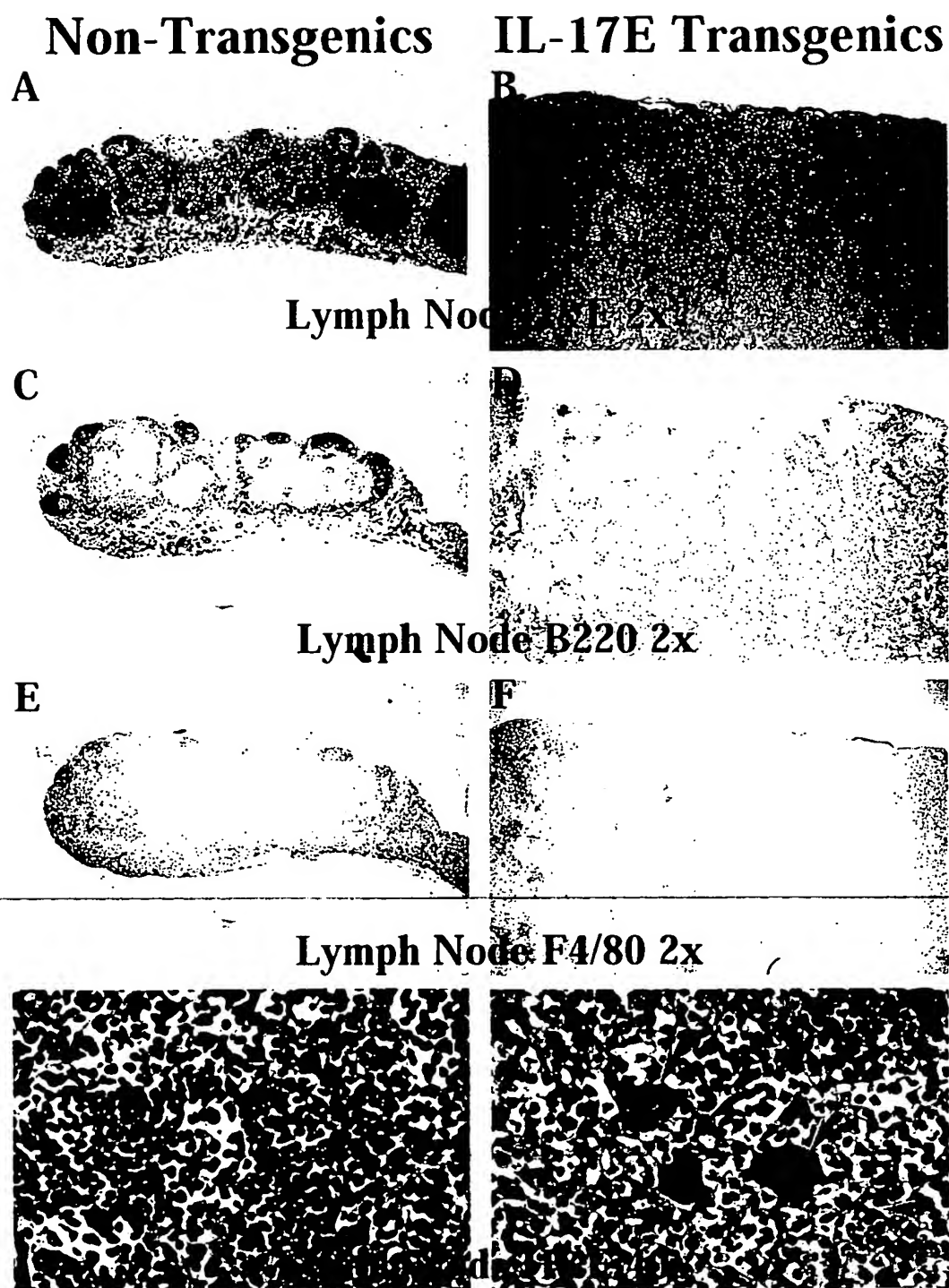
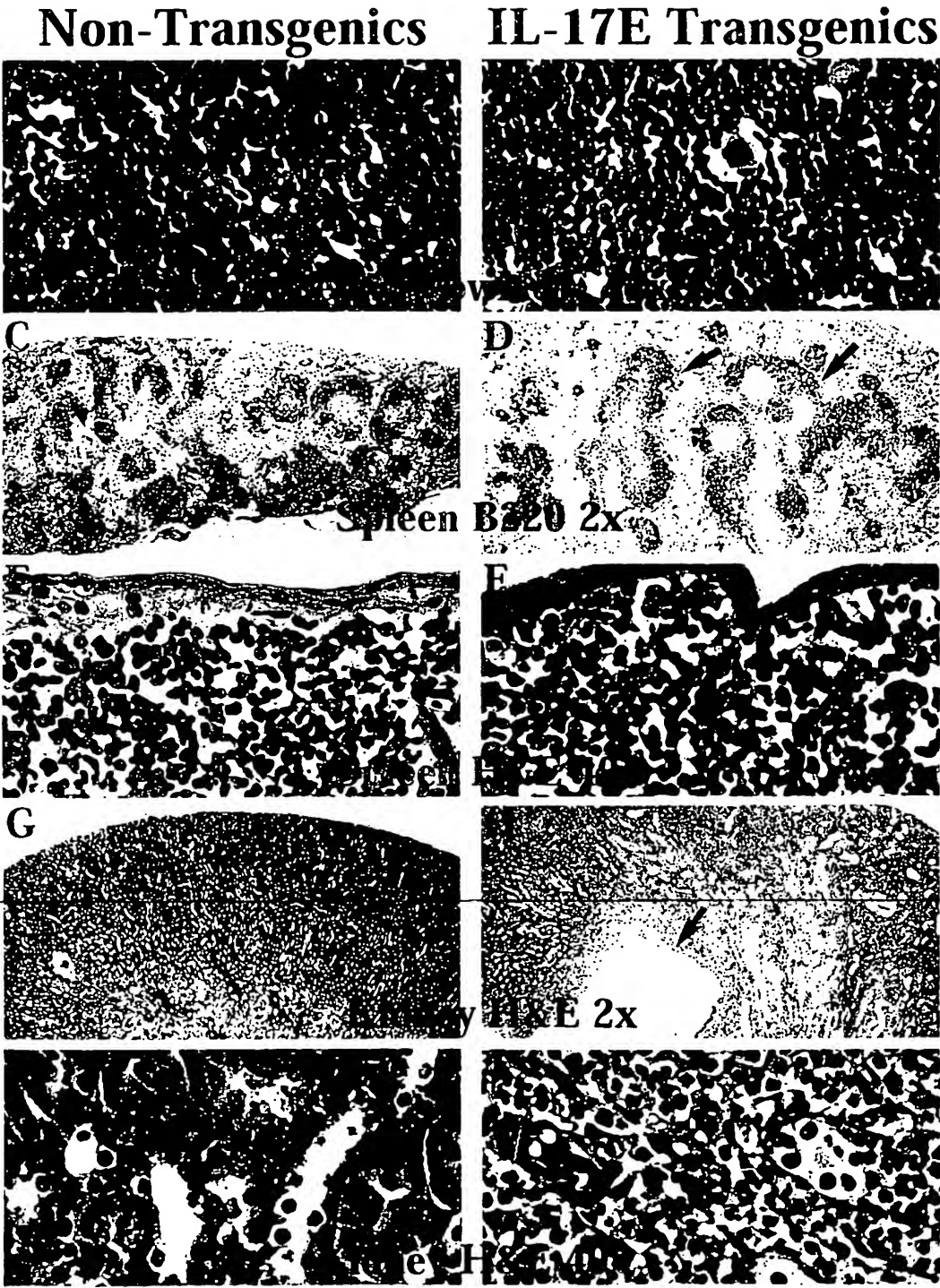


Figure 10



00010937 433304  
T05002 22607860

Figure 11



09340937 100004  
100004 100004

Figure 12

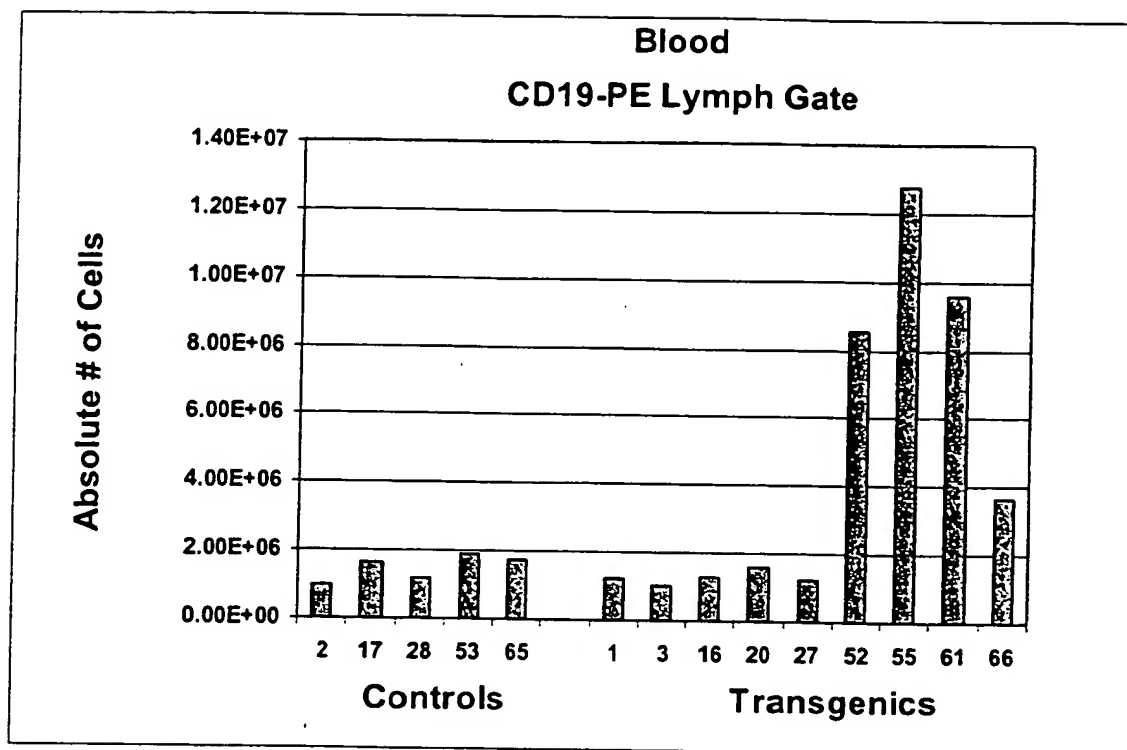




Figure 13

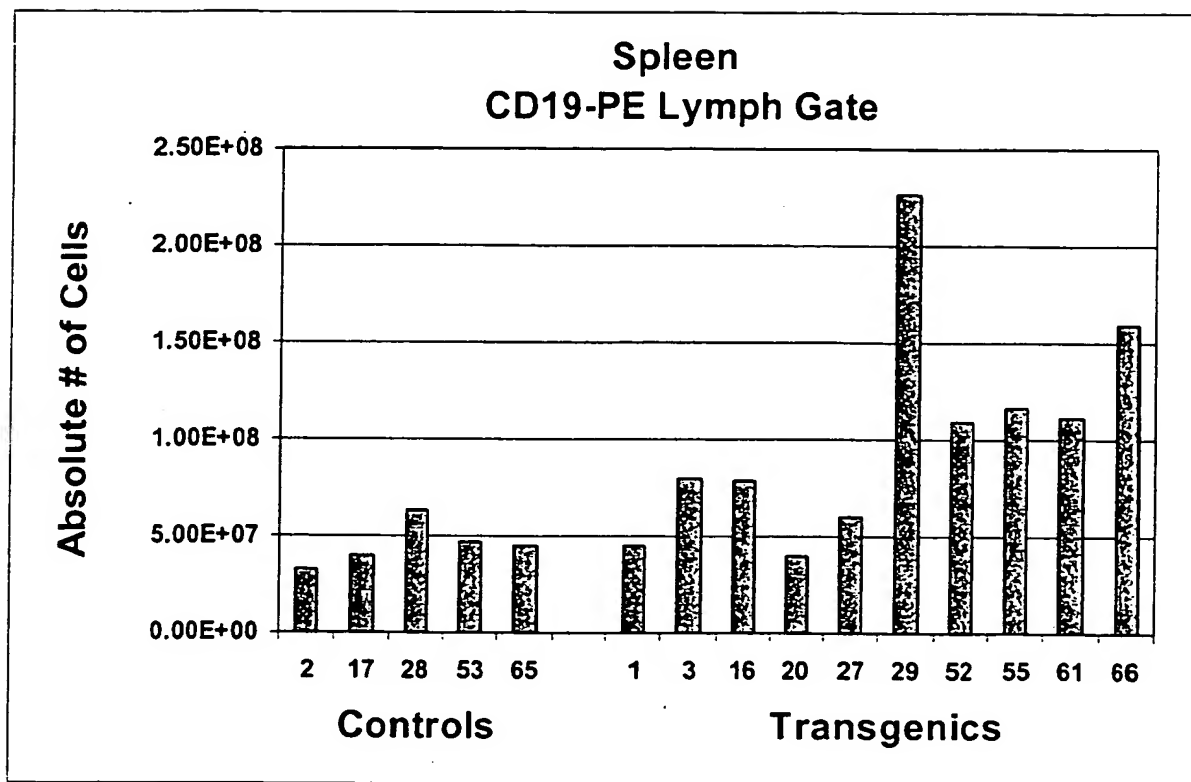


Figure 14

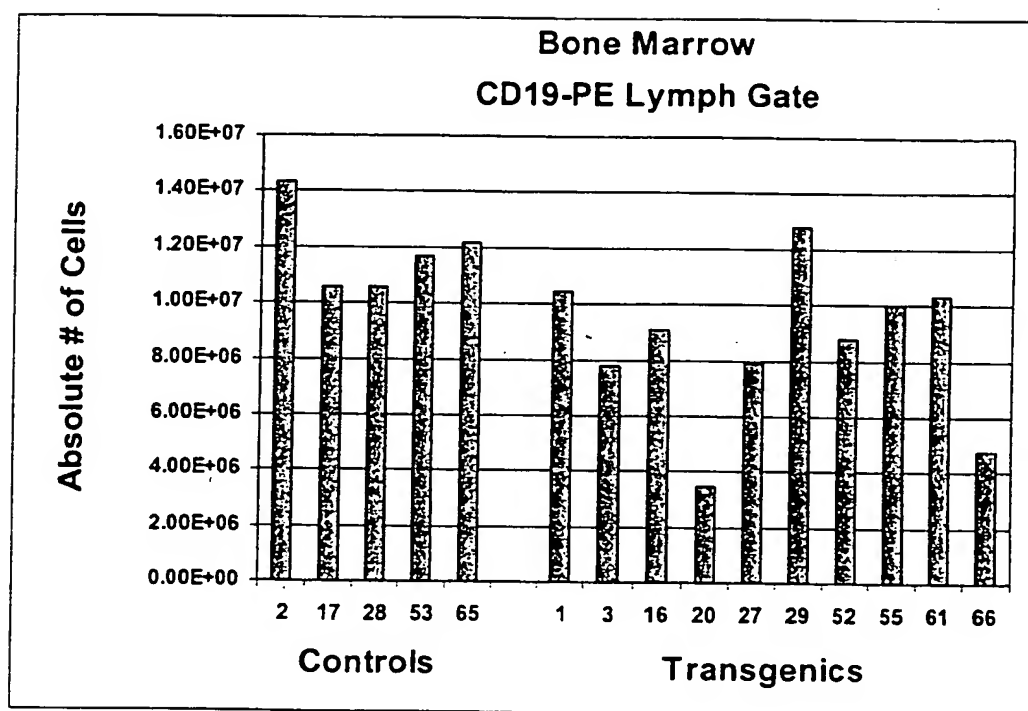


Figure 15

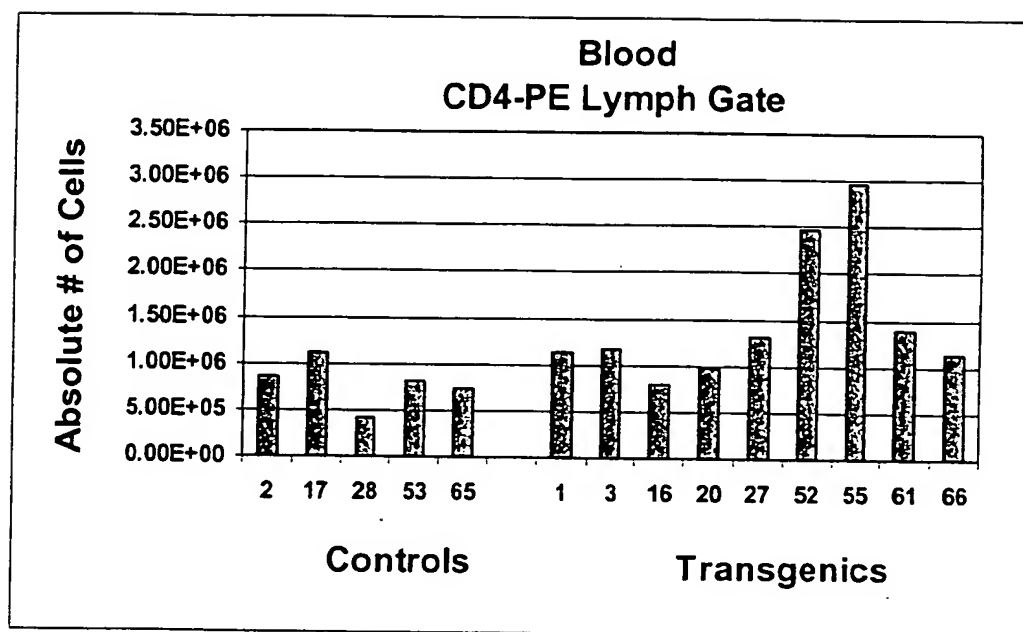


Figure 16

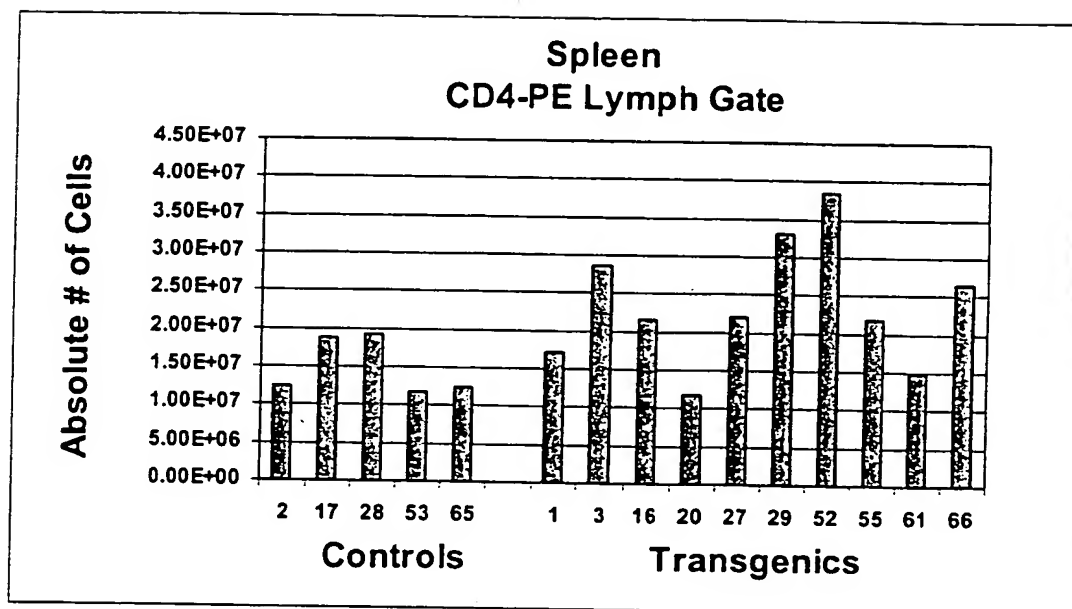


Figure 17

# CD45R+ CELLS EXPRESSING IL17Br IN TRANSGENIC BONE MARROW

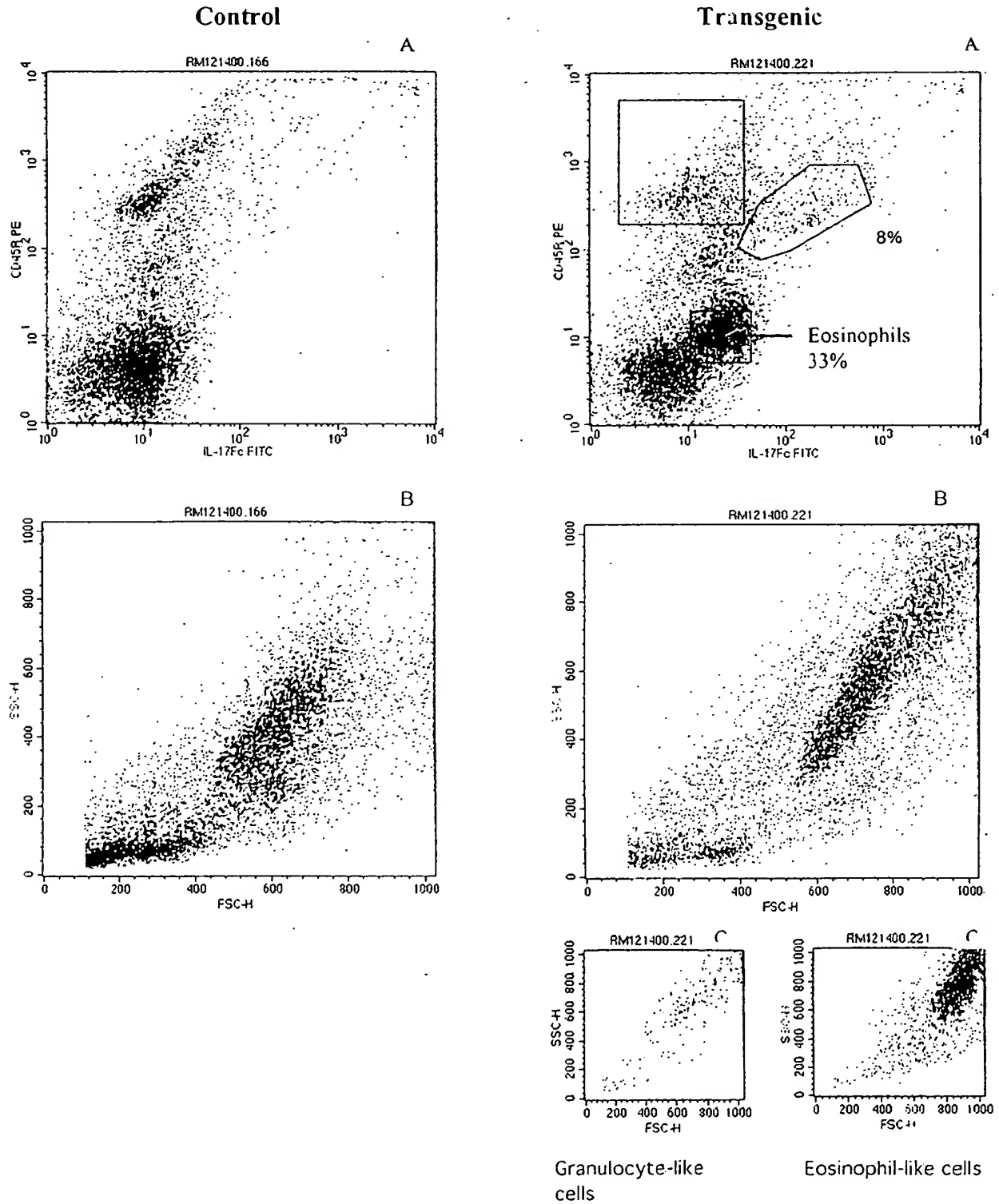
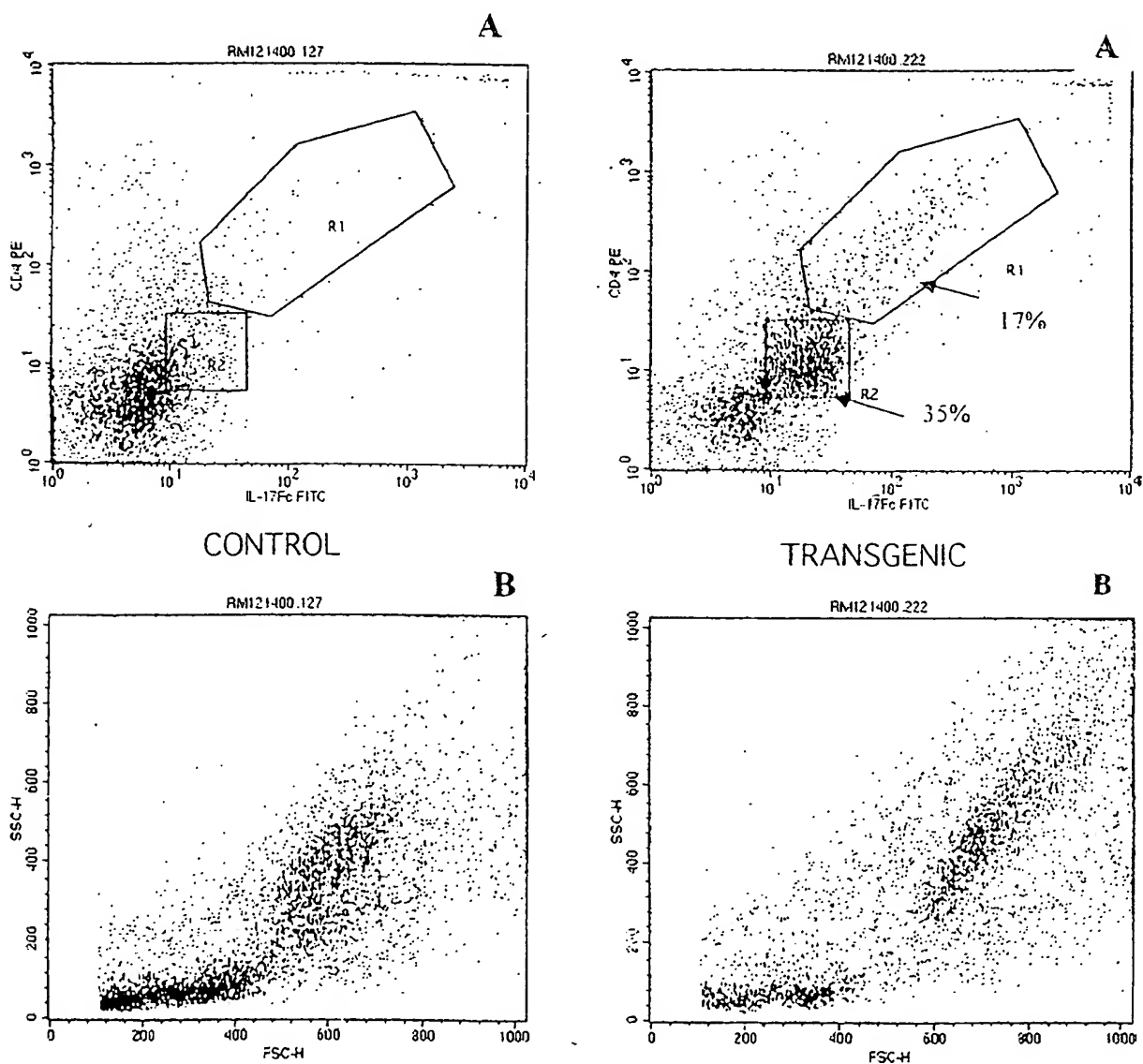


Figure 18

CD4+ CELLS EXPRESSING  
IL17Br IN TRANSGENIC BONE  
MARROW



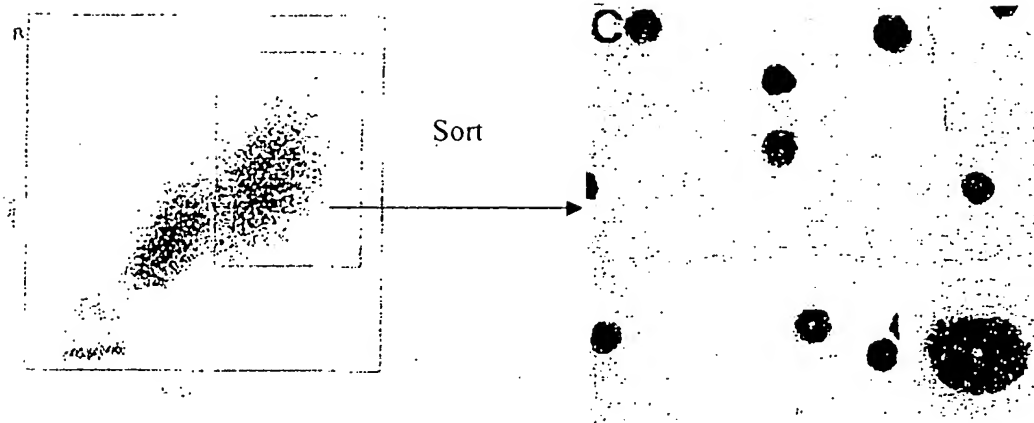
[illegible]

Figure 19

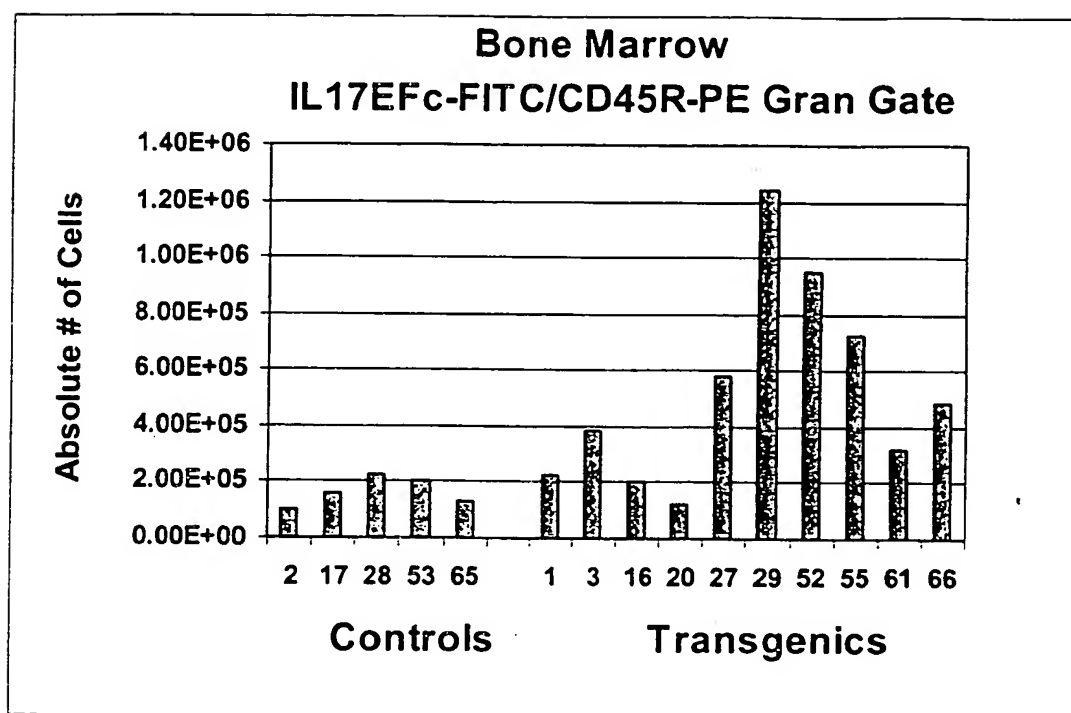
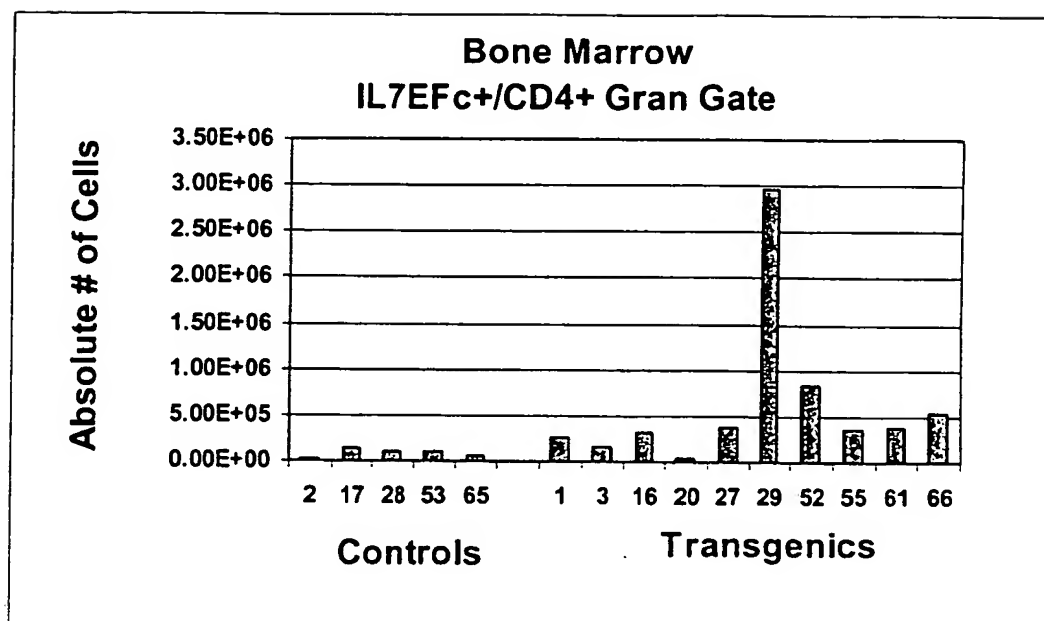




Figure 20



IL-17RB-2 Fusion Protein (SEQ ID NO: 24)

1	<u>MSLVLLSLAA</u>	<u>LCRS</u>	<u>AVPREP</u>	<u>TVQC</u>	<u>SETGP</u>	<u>SPEWMLQHD</u>	<u>LIPGDLRDLRV</u>
51	<u>EPVTTSVATG</u>	<u>DYSILMNVSW</u>	<u>VLRADASIRL</u>	<u>LKATKICVTG</u>	<u>KSNFQSYSCV</u>	<u>IPNANMNEDG</u>	<u>101</u>
101	<u>RCNYTEAFQT</u>	<u>QTRPSGGKWT</u>	<u>FSYIGFPVEL</u>	<u>NTVYFIGAHN</u>	<u>EETVEVNFTT</u>	<u>201</u>	<u>151</u>
151	<u>PSMSVNFTSP</u>	<u>GCLDHIMKYK</u>	<u>KKCVKAGSLW</u>	<u>DPNITACKKN</u>	<u>GDSEGATVQL</u>	<u>251</u>	<u>301</u>
201	<u>TPLGNRYMAL</u>	<u>IQHSTIIGFS</u>	<u>QVFEPHQKKQ</u>	<u>TRASVVIPT</u>	<u>LPAAAEPKSC</u>	<u>351</u>	<u>401</u>
251	<u>TPYFPTCGSD</u>	<u>CIRHKGTVVL</u>	<u>CPQTGVPFPL</u>	<u>DNNKSKPGGW</u>	<u>CVVVVDVSHED</u>	<u>451</u>	<u>501</u>
301	<u>DKTHTCPPCP</u>	<u>APELLGGPSV</u>	<u>FLFPPKPKDT</u>	<u>LMISRTPEVT</u>	<u>QDWLNGKEYK</u>	<u>501</u>	
351	<u>PEVKFNWYVD</u>	<u>GVEVHNAKTK</u>	<u>PREEQYNSTY</u>	<u>RVVSVLTVLH</u>	<u>NQVSLTCLVK</u>		
401	<u>CKVSNKALPA</u>	<u>PIEKTISKAK</u>	<u>GQPREPQVYT</u>	<u>LPPSRDELTK</u>	<u>TVDKSRWQQG</u>		
451	<u>GFYPSDIAVE</u>	<u>WESNGQPENN</u>	<u>YKTTTPVLDS</u>	<u>DGSFFLYSKL</u>			
501	<u>NVFSCSVMHE</u>	<u>ALHNHYTQKS</u>	<u>LSLSPGK*</u>				

[illegible]

Figure 23

Fusion Protein for IL-17RB-3 (SEQ ID NO: 25)

1    **MSLVLLSLAA** **LCRS**AVPREP TVQCGSETGP SPEWMLQHDL IPGDLRLDRV  
51   EPVTTTSVATG DYSILMNVSW VLRADASIRL LKATKICVTG KSNFQSYSCV  
101   RLECSGAIMA RCDLNLGSS DRSASASRAA GTAGVGHQTW LIFVVFVEGG  
151   FTVLLVLNSS AQAICLPRLP KVLGLQWTFS YIGFPVELNT VYFIGAHNIP  
201   NANMNEDGPS MSVNFTSPGC LDHIMKYKKK CVKAGSLWDP NITACKKNEE  
251   TVEVNFTTTP LGNRYMALIQ HSTIIGFSQV FEPHQKKQTR ASVVIPTGD  
301   SEGATVQLTP YFPTCGSDCI RHKGTVVLCP QTGVPFPLDN NKSKPGGWLP  
351   **AAAEPKSCDK** **THTCPPCPAP** **ELLGGPSVFL** **FPPKPKDTLM** **ISRTPEVTCV**  
401   **VVDVSHEDPE** **VKFNWYVDGV** **EVHNAKTKPR** **EEQYNSTYRV** **VSVLTVLHQD**  
451   **WLNGKEYKCK** **VSNKALPAPI** **EKTISKAKGQ** **PREPQVYTLF** **PSRDELTKNQ**  
501   **VSLTCLVKGF** **YPSDIAVEWE** **SNGQPENNYK** **TTPPVLDSDG** **SFFLYSKLTV**  
551   **DKSRWQQGNV** **FSCSVMHEAL** **HNHYTQKSLS** **LSPGK\***